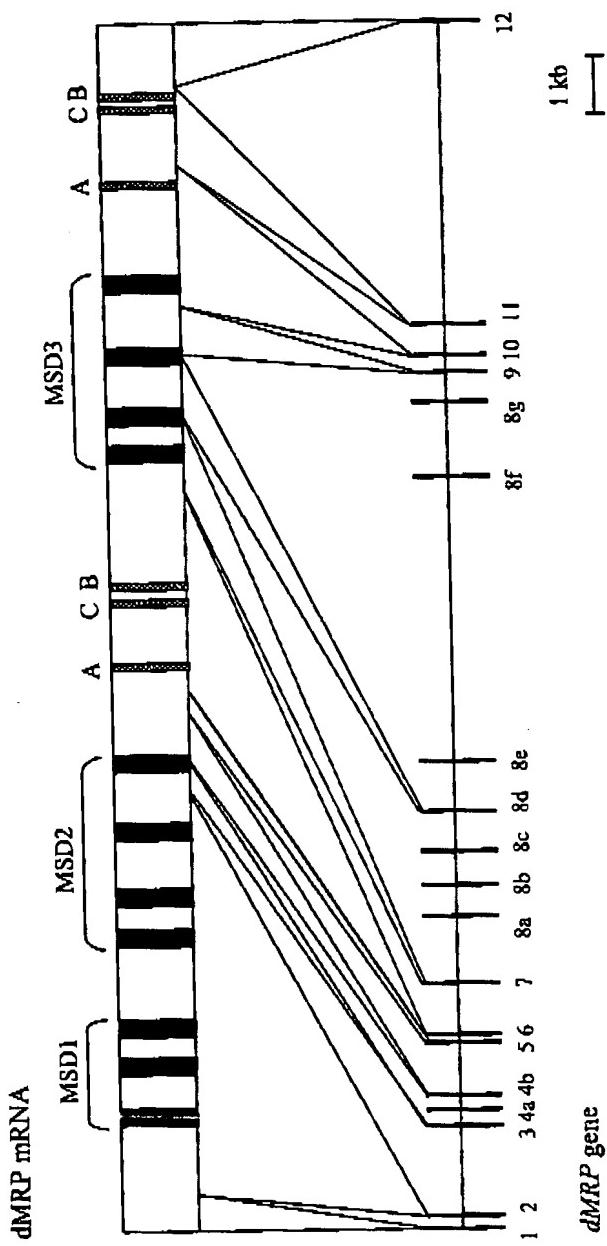


Figure 1

1 MADDTS P KDRPCGGSTFAN TUTS YN DPUETPCTEQTATYNGRPGAFYWAFTVIFDFYVLSKASIDORNIPWNYKXVSIALVNGLIUVITALDULMALKG-GDSEHPLPLYDLDVWCGPI
 HALRGPC-- SADGS DPLM DPNVWTNTSNPDFTKCFQNTVLUWVPCFYLNACPPFLYLSRNDRGYIQTNPPLNKTKTALCPFLWIVCHADLFYTSWERSRGTPLAFLVUS---PT
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Figure 2



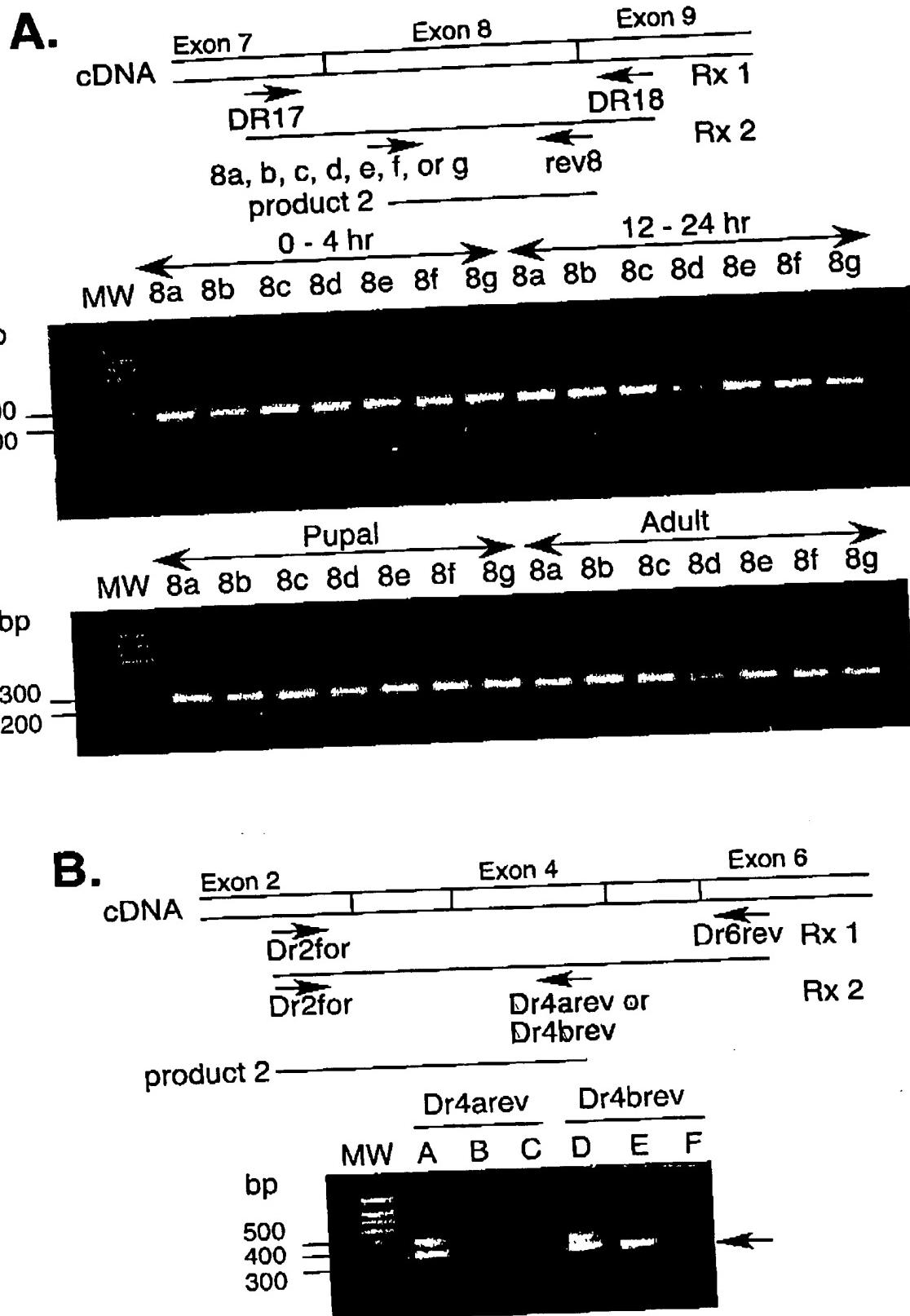


Figure 3

A	MRP1 MRP2 MRP3 4a 4b Dsur Cons	VALCCTFAYVVTIDENN--ILDAQTAFVSALAFNIRFPFLNILPVMVISSLVQ VSVVTTFSVVVLVDSDNN--ILDAQKAPTSITLNLIRFPPLSMLPMMISSMLQ VTLITLTMWVYVYDOPNN--VLDAAEKAFVSVSLFNILRRLPNNLPMOLISNLTQ VSLVLTPTAVLTLSEBN--QLSVEKTVRSLSALENIRFPFLNLPWILTNLVQ ITPVTTLGCVVWLNLRDQEFDLNASRFLSSSLAFQQLTVPFLLIFPITVPILIA V-LVTF-VVV--D-NN---LDA-K-FVS--LPNILR-PL--LPM-I---Q
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B	IAVFGYMAVSIIGGILASRCLEVDLHSLR-SPMSFFERTPSGNLVNRFSEKLDTVDShIPEVIKMFGSLEN IPVPIAHFWSAFEVHSNLHKLNNILR-APMRRPFDTTPTGRIVNNRAGDISTVDDTLPOSRLSWITCFGL FLVMLAAMAAAGGICQAARVLHQALEHNKR-SPQSFBDTTPSGRILNCFSRKDIYVVDLAVPLVNLLNSPN ATNNTTSSLAISLGCLQKCOLHQTTLYNNR-WPMENFDITPLGRIVNRFBSKDVTDTIDNTPLNLRVVLQLF TSYFFCSCITLALCCIPCSKVNLHETLSSVER-WPMELFDTTTPGRIVNNRFSEKSDVDTIDNVLPMLWRMVTOSAPA LSKYLSSLALGGLHCMSNNVNLNTGLK-WPMELFTTTPGRILNSPKDVTDTPLGRILNSPKDVTDTPLGRILNSPKD VLAYFAVIVVILVLCGPQAKATTIHNLAVIIRGSVCRPFEDITPGKRLNSPKDVTDTPLGRILNSPKDVTDTPLGRILNSPKD LCNYGAISLFTLTHASSRVFRHLNNIMH-CPSRFDFDTPLGRILNSPKDVTDTPLGRILNSPKDVTDTPLGRILNSPKD PTSRPSDLAPALGSLHAALKVLSMILENVLR-APMTMFDTTPVGRILSRESKDVESVUDOKMPQVINDCIWCAFE VSTPAGQYAGCNARLNLRDKL(TILKTKLH---FPOVNTPLGRILNSPKDVTDTPLGRILNSPKDVTDTPLGRILNSPKD LLQGSFYFGERLQQ-RLRKFRALRQ-LEMSFPLGRILNSPKDVTDTPLGRILNSPKDVTDTPLGRILNSPKD Dsur Pfam Cons
MRP1	8a
MRP2	8b
MRP3	8c
MRP4	8d
MRP5	8e
MRP6	8f
MRP7	8g
MRP8	Dsrr
MRP9	Pfam
MRP10	Cons

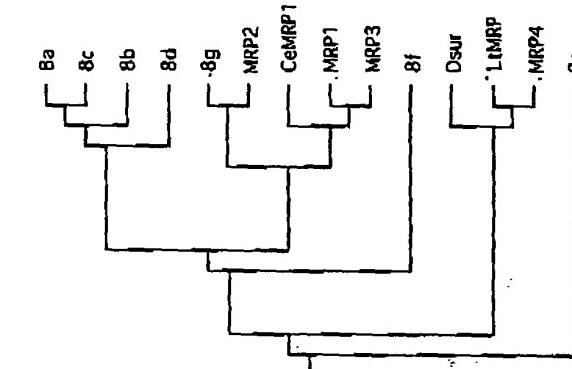
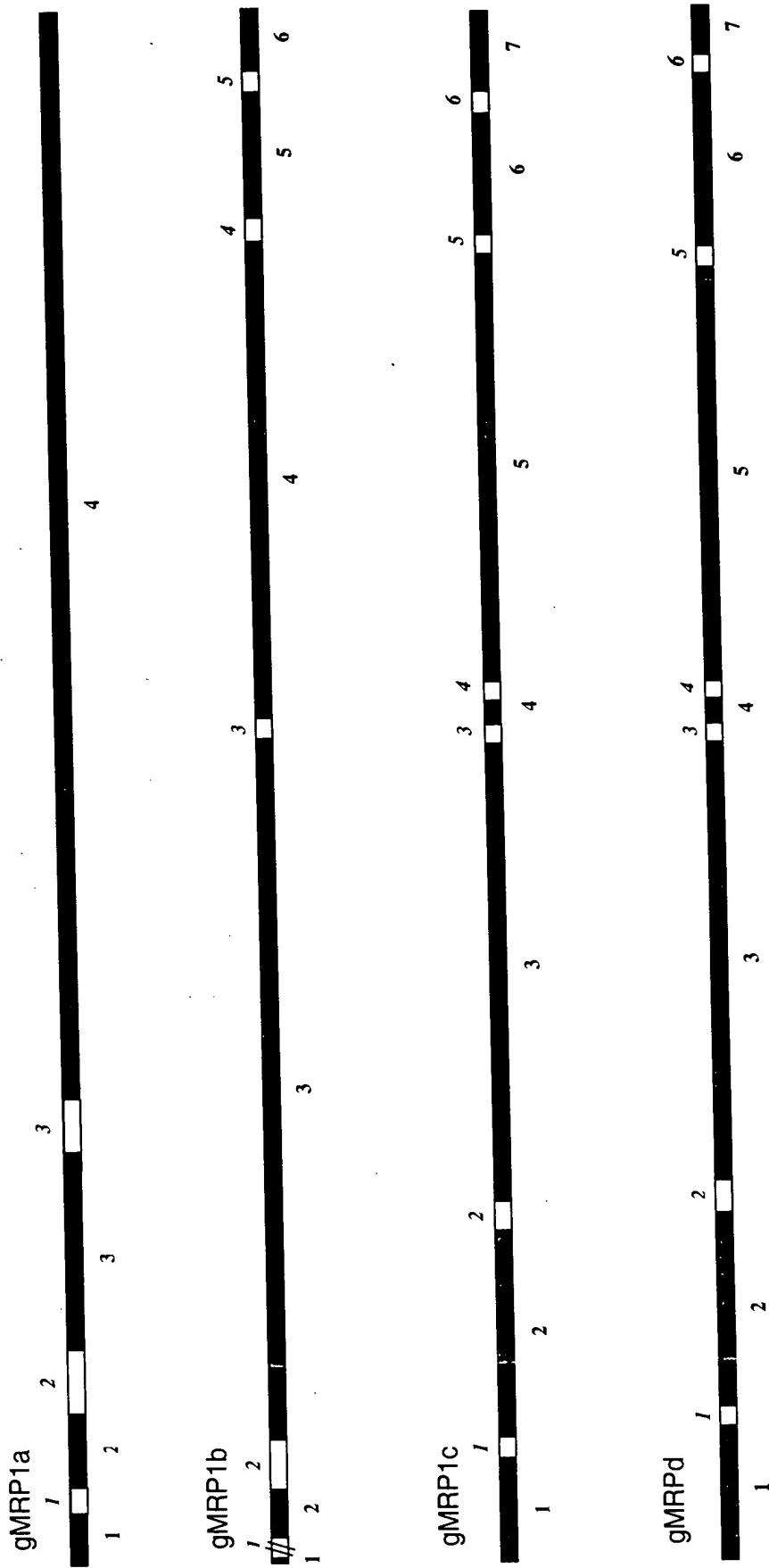


Figure 4

FIGURE 5



gMAP1c G F V W D N P N L T F C F Q R V I L Q W V P C L F L F V F S I Y D I 50
 gMAP1d P F W D G E F V W D N P N L T F C F Q R V I L Q W V P C L F L F V F S I Y D I 50
 gMAP1b P F W D D E F V W D N P N L T F C F Q R V I L Q W V P C L F L F V F S M Y E V 50
 gMAP1a P F W D D L T W R E E D P D L T F C F Q R V I L Q W V P C L F L F V Y L L V C L F F 27
 dMAP P V S L P L I V D H L V V Y G F Q Y L L V C L F F 27
 MAP1 M A D D T S S P M D R F C G S - - - T F W N A T E T W Y T N D P D F T S N P D F T K C F Q N T V L V W V P C F Y L W A C F P F Y F 54

 gMAP1c F K I T E S K Y R D I P W N W Y N L S K M L V I F L L M C M C W I D L G M V V G Y Q - - - D E Q G L Y D V Q I L T A V F 107
 gMAP1d F K I T E S K Y R D I P W N W Y N L S K M L V I F L L M C M C W I D L G M V V G Y Q - - - D E Q G L Y D V Q I L T A V F 107
 gMAP1b L R I V T S R Y R D I P W N W F N I T K M I F T F A L M V M S W V D L G V V V Q N L - - - D E P E V F D O V Q I L V A I F 107
 gMAP1a L S E L S S V K Q D V P V Q T S R A T V H S I F S A F L V L V T L A G V V V A A F R - - - L V D D S V A V V W R D G I 82
 dMAP L Y L K A S L D R N I P W N K L N V S S K A L V N L G L L V A M A L V K K G G D S E L D L D V W G P I I 116
 MAP1 L Y L S R H D R G Y I Q M T P L N K T K T A L G F L L W I V C W A D L F L Y S F W E R - - S R G I F L A P V F L V S P T L 112

MSD1

gMAP1c N A L A Y I I D L L V L L F M R K Y G V R T S G T M F M F W F L R M F F G I I Q L R A T E V M E N D K R P N A I G S G D - 166
 gMAP1d N A L A Y I I D L L V L L F M R K Y G V R T S G T M F M F W F L R M F F G I I Q L R A T E V M E N D K R P N A I G S G D - 166
 gMAP1b N A L A Y I I M A M A L Y F F Y R K Y G I R S T G T M F I F W F L K A F F G I I Q M R A T E V M E N D K R P N A I G S G D - 166
 gMAP1a E A I A S S L V I V G I L F L Q I Y S I R A N Y D I H I Y L F T F W T L R T L A L S M D V A F D A T - - - - - 161
 dMAP K F A T T I L L G I L F I F I P L N R K Y G V Q T T G C Q F I F W F L L T V L S I P R C R E V A R L D A E R Q K I L N S Q Q P 127
 MAP1 L G I T T I L L G I L F I F I P L N R K Y G V Q T T G C Q F I F W F L L T V L S I P R C R E V A R L D A E R Q K I L N S Q Q P 127

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gMAP1c - - - T V D F W E Y Q Y V S Y I L Q Y S L I C L M L V L E L F P D K E P S F S Y Y P K A A K P N P E L R S S F F A K L L 223
 gMAP1d - - - T V D F W E Y Q Y V S Y I L Q Y S L I C L M L V L E L F P D K E P T F S Y Y P K S K N R N P E L R S S F F A K L L 223
 gMAP1b - - - T G D F D A E F Q F V S Y T I Q Y T F V G C V L L E L F P D K E P R Y S E W A K L K N P N P E L R S S F F S R L F 218
 gMAP1a F S W M D S T Y R E A H R G S V A F Y Q G T L F Q Q T L F Q Q T L F Q Q T L F Q Q T L F Q Q T L F Q Q T L F 174
 dMAP S E Q D F S W E E Y Q F V S F T S I M L I L N C F A D G M P R Q T K Y Q R G N P E L S A S F L S R I T 226
 MAP1 - - - | D A Q V D L F R D I T F Y V Y F S L L I Q L V L S C F S D R S P L F S E T I H D P I N P C P E S S A S F L S R I T 226

MSD1

gMAP1c F L H F D A F A W K G F R N P L T M N D M Y D I N P Q D S A R E L V P P F D K Y W K I S V E K G R K - - - - - 274
 gMAP1d F L Y F D T F A W K G F R K P L T M E E M Y D I N P Q D T S R A L Y P P F D K Y W D M S V A N G R K - - - - - 274
 gMAP1b F L Y F D S Y A W R G F R K P L T D D M Y D L N P E D T S R A L Y P P F D K Y W E S V E K G R K - - - - - 269
 gMAP1a F S W M D S T Y R E A H R G S V A F Y Q G T L F Q Q T L F Q Q T L F Q Q T L F Q Q T L F Q Q T L F Q Q T L F 224
 dMAP Y Q W W I T G I L I V R G Y R Q P L E E G S D L W S L N K E D T S E Q V V P V L V K N W K K E C A K T R K I Q P V K V V Y S S K 297
 MAP1 F W W I T G I L I V R G Y R Q P L E E G S D L W S L N K E D T S E Q V V P V L V K N W K K E C A K T R K I Q P V K V V Y S S K 297

MSD2

gMAP1c Q M A S D R K A G K P D - - - - - D Y K - P H S P S N G - - - - - S V L Y T M I R A Y Q G P F W F A G M L Q L A 320
 gMAP1d Q I A A D K K A G K T N - - - - - I E Y K - P H S E T N G - - - - - S S L Y A M V R A Y G A P P F W F A G M L Q L A 320
 gMAP1b Q I A A D K K A G K T N - - - - - L V Y K - P N A A T N G - - - - - S V L P A M V K A Y G G P F W F A G M L Q F A 315
 gMAP1a Q I A A D K K A G K T N - - - - - F T I G K L L S P K F R G E I I L A G L N R F V 263
 dMAP K A R V E P K A Q Q F S N G - - - - - N V T F E N P H G E K N G A K K G M A S I M P P I Y K S F G G V F L F G G A L M K L F 342
 MAP1 D P A Q P K E S S K V D A N E E V E A L I V K S P Q K E W N P - - - - - S L F K V L Y K T F G P Y F L M S F F F K A I 334

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gMAP1c I S G L Q F A S P Y L M Q I E L M A V I A F D G - - P L W K G F L L T F G L F G A S L L L G L F N G Q Y L F Y T F L S Q F 378
 gMAP1d I S G L Q F A S P Y L M Q I E L M A V I A F D G - - P V V W K G L L L T F A L F A S S L L L G L F N G Q Y Y N T F L S Q F 378
 gMAP1b I S G L Q F A S P Y L M Q I E L M A V I A F D G - - P F W K G M I I T L G L F F D A S T I A I N T H Y Q H T D I G L 373
 gMAP1a L I S L L F C P Y L L R I L L I S F V E A Q D A E P E W K G I L Y A V L F V L A A Q T F I L G Q Y F H R M F I V G L 318
 dMAP T D T L T F A Q P Q V L S I K F V N D T K - A P D W Q G Y F Y T V L L F V T A C L Q T L V L H Q Y F H I C F V S G M 402
 MAP1 H D L M M F S G P Q I L K I L K F V N D T K - A P D W Q G Y F Y T V L L F V T A C L Q T L V L H Q Y F H I C F V S G M 393

Figure 6

9MRP1c	R I R T G L I S A I Y R K A L R I S S A A K K D O T T V G E I V N L M A V D A Q K F F E L L T S Y L H I L W S A L L I G U	438
9MRP1d	R I R T G L V S A I Y R K A L R I S S A A K K D O T T V G E I V N L M A V D A Q R F F E L L T S Y L H I L W S A P L I G U	438
9MRP1b	R I R T G L I S A I Y R K A L R I S S F A K K D O T T V G E I V N L M A V D A Q R F F E L L T S Y L H I L W S A P L I G U	433
9MRP1a	R I R S I L M G A I Y R S I L H D G I T S N A S S S D T I L T S D A Q R F F E L L T S Y L H I L W S A P L I G U	372
dMRP	R I R T A V I G A V Y R K A L V I T N S A R K S S T V G E I V N L M S V D A Q R F M D L A T Y I N M I W S A P L Q V I L	462
MRP1	R I R T A V I G A V Y R K A L V I T N S A R K S S T V G E I V N L M S V D A Q R F M D L A T Y I N M I W S A P L Q V I L	453
MSD2		
9MRP1c	C V F L L Y D - I L G P A V F A G L G Y M I L M T P V S G V V A A K L K T H Q V A Q M K L K D E R V K K M N E I L G G I	497
9MRP1d	C C V Y L L Y D - I L G P A V F A G L G Y M I L M T P V S G V V A A Q M K I K D D A Q Q V A Q M K I K D E R V K K M N E I L G G I	497
9MRP1b	C C I Y L L Y E - I L L G P A V F A G L G Y M V M I P I T G F I A T R M R D L Q V E Q M K I K D E R V K K M N E I L G G I	492
9MRP1a	T F V A L W V G V L G P I G T V G G L A I I V A V I A T R K L A K K I A A Q E K H I T A H S N D A V A L T T A S I E Q M	492
dMRP	A L Y F L W Q - Q L G P S V L A G G L A V M I L I P V N G V I A S R I K T Y Q I R Q M K Y K D E R V K L M N E V L S G I	432
MRP1	A L Y L L W L I - N L G P S V L A G G V A V M V L M V P V N A V V M A M K T K T Y Q V A H M M K S K D N R I K L M N E I L N G I	512
MSD2		
9MRP1c	K V L K L Y A W E P S F Q D S I L N V A R D E E V G I L K K M A Y Y G A G I F F T F T I A P F L V T L V S F A V Y V L I D	557
9MRP1d	K V L K L Y A W E P S F Q D N I L T V A R K E E I G I L K K M A Y Y G A G I Y F F T F T I A P F L V T L V S F A V Y V M I D	557
9MRP1b	K V L K L Y A W E P S F Q D T V V T V R N E E L D V L K S A A Y Y G A G T Y F V W T M A P F L V T L V S F A V Y V M I D	552
9MRP1a	Q Q I K S D L M E P P F E Q R A I G E H R R A E L T H M C T Y I L Y D A L K Y L L S I A T P M I V A C G T F L F M Y V V G	492
dMRP	K V L K L Y A W E P S F E K Q V L D I R D K E I A T L L R S T A Y L N A D G T S F L W S C A P F L I V S L V T F A V Y V L I D	581
MRP1	K V L K L Y A W E L A F K D K Y L A I R Q E E L K V L K K S A Y L S A V G T F T W V C T P F F L I V A L C T F A V Y V T I D	572
MSD2		
9MRP1c	E N N V L D P Q T A F V S L A L F N I M R F P L G M F P M V V T F S M Q A W V S V K R A I D K F L N S A E L D P N N - -	614
9MRP1d	E E N V L D P Q T A F V S L A L F N I L R F P L G M L P M M M I T F A M Q A W V S V K R A I D K F L N S A E L D P N N - -	614
9MRP1b	P O N G S S T G G A G R G S F E K M Q E V V H T F V D Q L E D S I A D T S R A E V R I E R A K F S T E K N T I L R G I	609
9MRP1a	S G A L L T V Q S M F V A I A L F G L T R Y P L S E L P N L M T N W G T I N V K L Q V I N E V V C S G K Q R K S S G K M	552
dMRP	E N N V L D A T K T F V S L S L F N I L R F P L T M L P M L I T N L V Q I T Q V S V N R I N K F L N S E E L D P N S - -	638
MRP1	E N N V L D A Q T A F V S L A L F N I L R F P L N I L P M V I S S I V Q I A S V S V L K R L A I F L S H E E L E P D S - -	629
MSD2		
9MRP1c	- -	643
9MRP1d	- -	643
9MRP1b	- -	638
9MRP1a	- -	612
dMRP	- -	667
MRP1	- -	684
NBD1		
9MRP1c	N L S L R K G Q L S A I V G T V G T G K S S S L I S A L L G E M E K I S G H V N T D G - S I A Y V P Q Q A W I Q N A T L R	702
9MRP1d	N L L A L R K G Q L S A I V G T V G T G K S S S L I S A L L G E M E K I M K G S S V N T D G - S I A Y V P Q Q A W I Q N A T L R	702
9MRP1b	N L T L R E G T F I G V S G T H G S S G K T S S L I S A L L G E M E K I G R L Q R T G G T S V N T D G - S I A Y V P Q Q A W I Q N A T L R	687
9MRP1a	N I E V K K G S L V A L V G T V G S S G K T S S L I S A L L G E M E K I L A G V V N T V G - K L A Y Y V P Q Q A W I Q N A T L R	672
dMRP	T F S I P E G A L V A V V G Q V G C G K S S S L I S A L L G E M E K I A E M D K V E G H V A I K G - S V A Y Y V P Q Q A W I Q N A T L R	726
MRP1	T F S I P E G A L V A V V G Q V G C G K S S S L I S A L L G E M E K I A E M D K V E G H V A I K G - S V A Y Y V P Q Q A W I Q N A T L R	723
NBD1		
9MRP1c	D N I L F G K A F D Q R K Y D N V I E C C A L R P D L E M L P G G D S T E I G E K G I N L S G G Q K Q R V A L A R A V Y	762
9MRP1d	D N I L F G K A F D Q R K Y D N V I E C C A L R P D L E M L P G G D S T E I G E K G I N L S G G Q K Q R V A L A R A V Y	762
9MRP1b	D N I L F G P F D Q Q A K Y D K V I E C C A L R P D L E M L P G G D T T E I G E K G I N L S G G Q K Q R V A L A R A V Y	757
9MRP1a	S N I L F G Q Q T Y D R K R Y N K V I D A C A L R A D I D I L S A Q D L T E I G E K G I N L S G G Q H S L S G G Q A R R V S L A R A V Y	732
dMRP	D N I L F G C Q L E E P Y Y R A S V I Q A C A L L P D L E I L P S Q D R T E I G E K G I N L S G G Q K Q R V S L A R A V Y	786
MRP1	E N I L F G V N L S G G Q K Q R V S L A R A V Y	783

Figure 6

3MAP1c	ADAEEVYLFDODPLSSAVDAHVGGKHI	FEKVIG	G - PSSGMLVGSRLLVTHGISSYLPFVENIFVVK	821	
3MAP1d	ADSEEVYLFDODPLSSAVDAHVGGKHI	FEKVIG	G - PSSGMLVGSRLLVTHGISSYLPFVENIFVVK	821	
3MAP1b	AHDADLYLDDPLSSAVDAHVGGKHI	FEEVVIG	G - PQHQLLAGCTVFLSHDPELKAQSVLADKVLVMA	816	
3MAP18	SNADEIYLDDPLSSAVDAHVGGKHI	FENVIG	G - PKGMELKNSMSSYLPQVDVIMSV	792	
3MAP1	DGEIISESGTQQIELLDQKGAFAAEFLSQHQELDQKGAFAAEFLTQH	QELDDEDE	IEISLIQETLND - - - - - GVVN	845	
3MAP10	DGEVSESGSYQIELLDQKGAFAAEFLTQH	QELDDEDE	DIEIKLQEAALKD - - - - - ETAQ	842	
3MAP1b	GGG - - - TIEKV	QELDDEDE	DELKLQEAALKD - - - - - GEAK	888	
3MAP18	MGGEISEMGSYQIELLDQKGAFADFLQH	QELDDEDE	EEKAQADQKQNSSTADVPELLGTV	894	
3MAP	GGK - - - TIEKV	QELDDEDE	EELNQVTGVSGPG - - - - - KEA	904	
3MAP1	DGEIISESGTQQIELLDQKGAFAAEFLSQHQELDQKGAFAAEFLTQH	QELDDEDE	ENIQVTGVSGPG - - - - - KEA	883	
Linker					
3MAP1c	NVIQRA - - - - -	VERS - - - - -	LSVRSNRSSNGSDGSTRKKP - - - - - VHS - KTVTTVPGRANL	921	
3MAP1d	KIVQRA - - - - -	MSTRSQRSGSSNGS	LSVRSGRSSGGSSNGS - - - - - AKKEDPPIQNLDKATL	923	
3MAP1b	- - - - -	MSTRSQRSGSSNGS	LSVRSGRSSGGSSNGS - - - - - QESKAS - AKEPDPIQNLDKATL	920	
3MAP18	- - - - -	MSTRSQRSGSSNGS	LSVRSGRSSGGSSNGS - - - - - SAEURNSKPRAVEQTVAQQQSSATL	889	
3MAP	- - - - -	MSTRSQRSGSSNGS	LSVRSGRSSGGSSNGS - - - - - KARAKAQ - KARAKAQ	841	
3MAP1	- - - - -	MSTRSQRSGSSNGS	LSVRSGRSSGGSSNGS - - - - - QDSHDS - VASAASLKKQEEETWKL	983	
3MAP10	- - - - -	MSTRSQRSGSSNGS	LSVRSGRSSGGSSNGS - - - - - MSD3 - VASAASLKKQEEETWKL	947	
Linker					
3MAP1c	IGVEEESATGAVTWL	VYKKYIQS	IGFKFGFGGSVLFSTA - - - - - INQGSGIYSSMWLTDWSEEDPDAAT	981	
3MAP1d	IGVEEESATGAVTWL	VYKKYIQS	IGFKFGFGGSVLFSTA - - - - - INQGSGIYSSMWLTDWSEEDPDAAT	983	
3MAP1b	IGVEEESATGAVTWL	VYKKYIQS	IGFKFGFGGSVLFSTA - - - - - INQGSGIYSSMWLTDWSEEDPDAAT	980	
3MAP18	IGVEEESATGAVTWL	VYKKYIQS	IGFKFGFGGSVLFSTA - - - - - INQGSGIYSSMWLTDWSEEDPDAAT	901	
3MAP	IGVEEESATGAVTWL	VYKKYIQS	IGFKFGFGGSVLFSTA - - - - - INQGSGIYSSMWLTDWSEEDPDAAT	1023	
3MAP1	IGVEEESATGAVTWL	VYKKYIQS	IGFKFGFGGSVLFSTA - - - - - INQGSGIYSSMWLTDWSEEDPDAAT	1007	
Linker					
3MAP1c	DPSVARDKYLGVYGA	LGAAQSI	DPSVARDMYLGVYGAQSI - - - - - KAAKESHNKLLESCLRAMPMSFFFDTT	1041	
3MAP1d	DPSVARDMYLGVYGA	LGAAQSI	DPSVARDMYLGVYGAQSI - - - - - KAAKESHNKLLESCLRAMPMSFFFDTT	1043	
3MAP1b	DPSVARDMYLGVYGA	LGAAQSI	DPSVARDMYLGVYGAQSI - - - - - KAAKESHNKLLESCLRAMPMSFFFDTT	1040	
3MAP18	GALLHTTWILC	VYGVVGL - - - - -	GALLHTTWILC	KAARELHSNLLLESCLRAMPMSFFFDTT	955
3MAP	DGTGLRDMDYLGVYGA	QIVLSKYLSQGLALA	DGTGLRDMDYLGVYGAQIVLSKYLSQGLALA	MSD3 - FDLN	1089
3MAP1	TQEHTKVRLS	VYGGALQISQGI	TQEHTKVRLS	MSD3 - FDLN	1067
MEADKAQ	MEADKAQ	VYWDYMKAVG	MEADKAQ	MSD3 - FDLN	1067
Linker					
3MAP1c	PLGRIINAFSKDV	DVVDNNVL	PLGRIINAFSKDV	PLGRIINAFSKDV	1104
3MAP1d	PLGRIINAFSKDV	DVVDNNVL	PLGRIINAFSKDV	PLGRIINAFSKDV	1103
3MAP1b	PLGRIINAFSKDV	DVVDNNVL	PLGRIINAFSKDV	PLGRIINAFSKDV	1100
3MAP18	PLGRIINAFSKDV	DVVDNNVL	PLGRIINAFSKDV	PLGRIINAFSKDV	1015
3MAP	PLGRIINAFSKDV	DVVDNNVL	PLGRIINAFSKDV	PLGRIINAFSKDV	1143
3MAP1	PLGRIINAFSKDV	DVVDNNVL	PLGRIINAFSKDV	PLGRIINAFSKDV	1122
Linker					
3MAP1c	FVQRA - - - - -	FVQRA - - - - -	FVQRA - - - - -	1154	
3MAP1d	FVQRA - - - - -	FVQRA - - - - -	FVQRA - - - - -	1155	
3MAP1b	LLIQK - - - - -	LLIQK - - - - -	LLIQK - - - - - FVQRA - - - - -	1153	
3MAP18	MLVQCG - - - - -	MLVQCG - - - - -	MLVQCG - - - - - LLQYNETQG - - - - - LLQYNETQG - - - - - LLQYNETQG - - - - -	1055	
3MAP	FAQQR - - - - -	FAQQR - - - - -	FAQQR - - - - -	1198	
3MAP1	FVQIR - - - - -	FVQIR - - - - -	FVQIR - - - - -	1180	

Figure 6

gMAP1c	RVDYNOVLVSYPT	VANRWLAVERLELIGS	CVIILFAALFAILARDT	GQATVGSISYALQI	I	1214
gMAP1d	RVDYNOVTYPSS	VANRWLAVERLELIVGSL	VIFALFAMVARDT	GQATVGLSISYALQI	I	1216
gMAP1b	RVDYNOQLTYPSS	VANRWLAVERLELIVGSL	VIFALFAMVARDT	GQATVGLSISYALQI	I	1213
gMAP18	RVDYDTHQNYIYHN	VANRWLAVERLELIVGAI	VIFALFAMVARDT	GQATVGLSISYALQI	I	1135
dMRP	KVDKNQVCKYPS	VANRWLAVERLELIVGAI	VIFALFAMVARDT	GQATVGLSISYALQI	I	1255
MAP1	KVDENQKAYYPSI	VANRWLAVERLELIVGAI	VIFALFAMVARDT	GQATVGLSISYALQI	I	1240
gMAP1c	SHYLSFLVVRMTSE	VAVELLEETNVAVAEWQKG	-	-	-	1289
gMAP1d	SNVLSFLVVRMTSE	VAVELLEETNVAVAEWQKG	-	-	-	1271
gMAP1b	SATLSSFLVVRMTSE	VAVELLEETNVAVAEWQKG	-	-	-	1268
gMAP18	IPSLNSLLVVRMSSD	VAVELLEETNVAVAEWQKG	-	-	-	1185
dMRP	TQTLNWLVVRMSSD	VAVELLEETNVAVAEWQKG	-	-	-	1313
MAP1	TTYNWLVVRMSSD	VAVELLEETNVAVAEWQKG	-	-	-	1297
<i>NBD2</i>						
gMAP1c	YQIARYREGLDLV	LNVRGGEEKIGI	VGRTGAGKSSSLTLLG	VEAAG-GQIIDGL	I	1328
gMAP1d	YQIARYREGLDLV	LNVRGGEEKIGI	VGRTGAGKSSSLTLLG	VEAAG-GQIIDGL	I	1330
gMAP1b	FSLTHADGG-DLVL	LNVRGGEEKIGI	VGRTGAGKSSSLTLLG	VEAAG-GQIIDGL	I	1327
gMAP18	FQVRYREGLDLV	LNVRGGEEKIGI	VGRTGAGKSSSLTLLG	VEAAG-GQIIDGL	I	1254
dMRP	YCLRYREDLDLV	LNVRGGEEKIGI	VGRTGAGKSSSLTLLG	VEAAG-GQIIDGL	I	1372
MAP1	YCLRYREDLDLV	LNVRGGEEKIGI	VGRTGAGKSSSLTLLG	VEAAG-GQIIDGL	I	1356
gMAP1c	DISKMGLHQALRGRL	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1388
gMAP1d	DISKMGLHQALRGRL	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1380
gMAP1b	DISKMGLHQALRGRL	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1387
gMAP18	DISMGRISLQKLRLGE	IPQSTSSLFSGGVQQN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1307
dMRP	DIAASMGLHMLASRAL	IPQDPVLFSGSLRMIN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1432
MAP1	DIAKIGLHDLRFKITI	IPQDPVLFSGSLRMIN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1316
<i>B</i>						
gMAP1c	AGLDHEIAENGGENL	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1448
gMAP1d	AGLDHEIAENGGENL	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1450
gMAP1b	AGLDHEIAENGGENL	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1447
gMAP18	AGLDHEIAENGGENL	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1364
dMRP	AGLDHEIAENGGENL	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1492
MAP1	AGLDHEIAENGGENL	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1476
<i>C</i>						
gMAP1c	AGLDHEIAHRLNTI	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1505
gMAP1d	AGLDHEIAHRLNTI	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1507
gMAP1b	AGLDHEIAHRLNTI	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1504
gMAP18	AGLDHEIAHRLNTI	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1414
dMRP	AGLDHEIAHRLNTI	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1548
MAP1	AGLDHEIAHRLNTI	IPQDPVLFSGTLRAN	FVKGGLSDQVWKA	VEAAG-GQIIDGL	I	1531
<i>NBD2</i>						
gMAP1c	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	A	1505
gMAP1d	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	A	1507
gMAP1b	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	A	1504
gMAP18	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	A	1414
dMRP	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	A	1548
MAP1	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	ADCTILTIAHRLNTI	A	1531

Figure 6